

FIG. 1

Intact Double Stranded DNA Chemical Fragmentation Randomly Fragmented Single Stranded DNA RHP Fill In Reaction Adaptor Ligation Extension **X**—

FIG. 2

Intact Double Stranded DNA Chemical Fragmentation Randomly Fragmented Single Stranded DNA RHP Fill In Reaction Ligation Adaptor Extension

FIG. 3

Double Stranded DNA

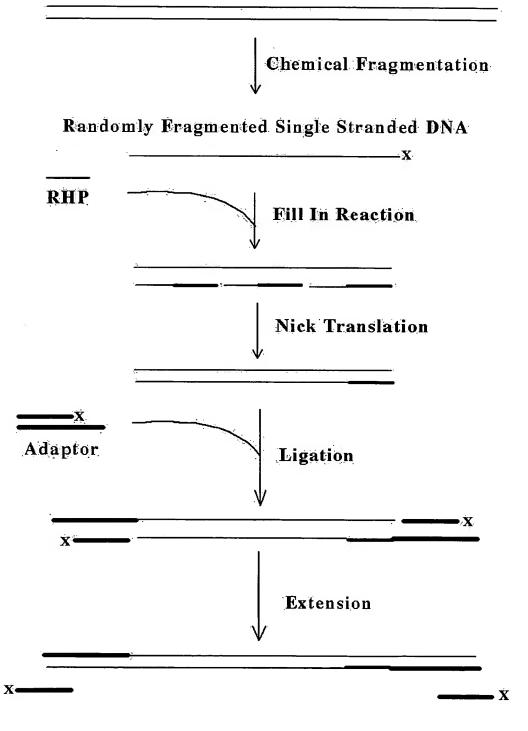


FIG. 4

Α

Blunt End Adaptor

5'GTAATACGACTCACTATAGG-3' (SEQ ID NO:32) 3'.XTGACGTGATATCC-5" (SEQ ID NO:33)

3' Overhang Adaptor

5'GTAATACGACTCACTATAGGN-3' (SEQ ID NO:34) 3'-XTGACGTGATATCC-5' (SEQ ID NO:33)

5' Overhang Adaptor

5'GTAATACGACTCACTATAGG-3' (SEQ ID NO:32) 3'-XTGACGTGATATCCN-5' (SEQ ID NO:35)

В

Linear T7HEG Adaptor

5'-CCTATAGTGAGTCGTATTACTTT HEG TTTTGTAATACGACTCACTATAGG-3'(SEQ ID NO:36)

Annealed T7HEG Adaptor

GTAATACGACTCACTATAGG-3'
CATTATGCTGAGTGATATCC-5' (SEQ ID NO:36)

S'-GTAATACGACTCAGTATAGG-3'(SEQ ID NO:32) 3'QTGAGTGATATCC (SEQ ID NO:33)	DNA Ligase	ATACGACTCACTATAGG COTATACTGAGT®3 (SEQID NO:33) (3) 3'-®TGAGTGATATCC GGATATCACTGAGCATATG-5' (SEQID NO:32)	Taq Polymerase	AATACGACTCACTATAGG COTATAGTGAGTCGTATTAC.3; (SEQ ID NO:37 (SATATGCTGAGTCGTATTAC.3; (SEQ ID NO:37 (SATATGCTGAGTGATTAC.5; (SEQ ID NO:32 (SATATGCTGAGTGATTG-5; (SEQ ID NO:32 (SATATGCTGAGTGATTGCTGAGTGTG-5; (SEQ ID NO:32 (SATATGCTGAGTGATTGCTGTGAGTGCTGATTGCTGTGTGTG
		(SEQID NO:32) S'GTAATACGACTCACTATAGG- (SEQID NO:33) S'GTGAGTGATATCC		(SEQ ID NO:32) S: GTAATACGACTCACTATAGG — (SEQ ID NO:37) 3'-CATTATGCTGAGTGATATGC —

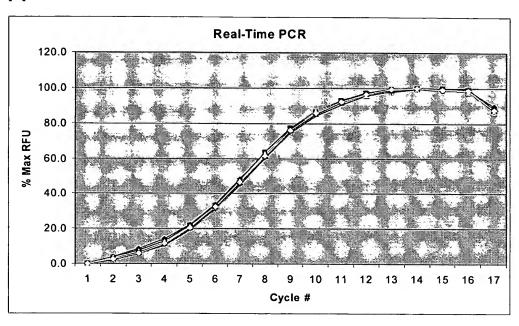
Fragmented DNA

S-CCTATACTGAGT®3" (SEQ ID NO:33)

denotes 3' blocking group

FIG. 6

Α



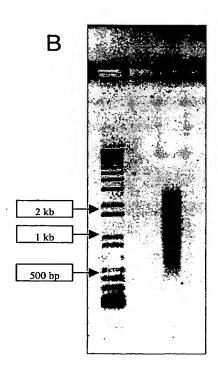
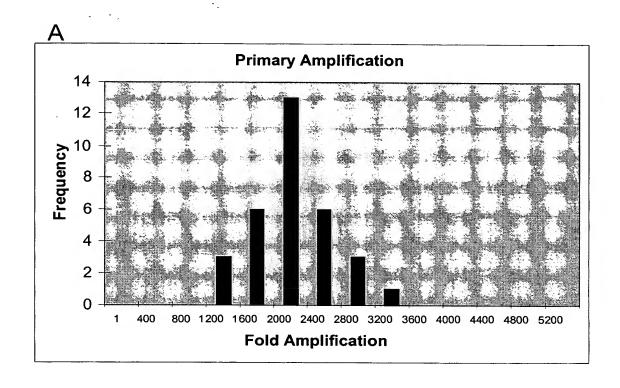


FIG. 7



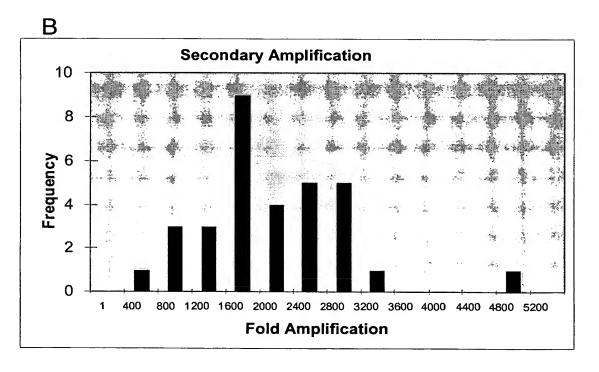
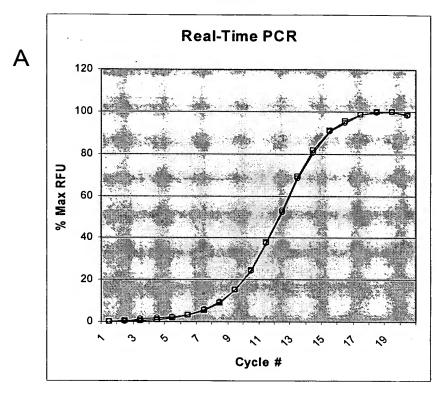


FIG. 8



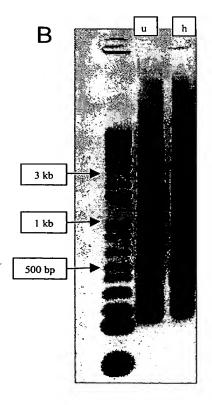


FIG. 9

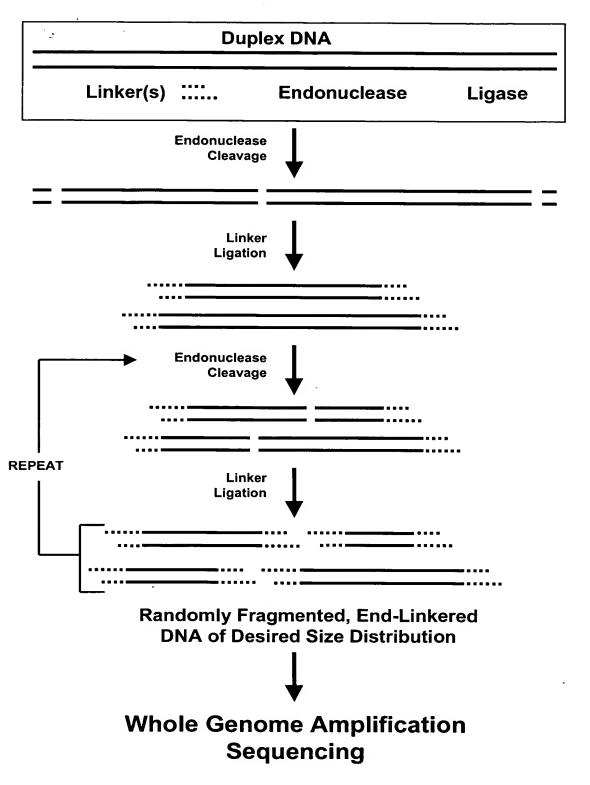


FIG. 10

11A	5' 3' 3' 5'
11B	5' N 3' 3'
11C	5' 3' 3' N 5'

FIG. 11

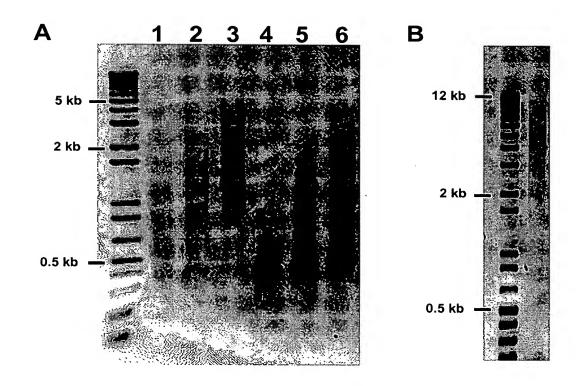


FIG. 12

```
A 5'-CCTATAGTGAGTX-3' (SEQ ID NO:33)
3'-GGATATCACTCAGCATAATGA-5' (SEQ ID NO:38)

B 5'-NCCTATAGTGAGTX-3' (SEQ ID NO:39)
3'-GGATATCACTCAGCATAATGA-5' (SEQ ID NO:38)

C 5'-NNCCTATAGTGAGTX-3' (SEQ ID NO:39)
3'-GGATATCACTCAGCATAATGA-5' (SEQ ID NO:38)

D 5'-CCTATAGTGAGTX-3' (SEQ ID NO:33)
3'-NNGGATATCACTCAGCATAATGA-5' (SEQ ID NO:40)

E 5'-CCTATAGTGAGTX-3' (SEQ ID NO:33)
3'-NGGATATCACTCAGCATAATGA-5' (SEQ ID NO:41)
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FIG. 13

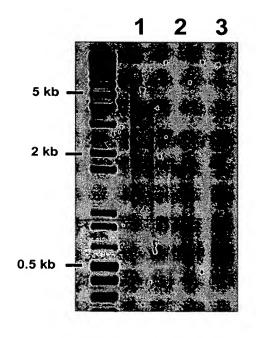
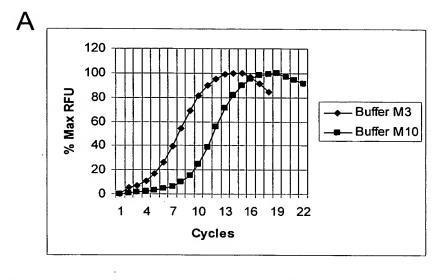


FIG. 14



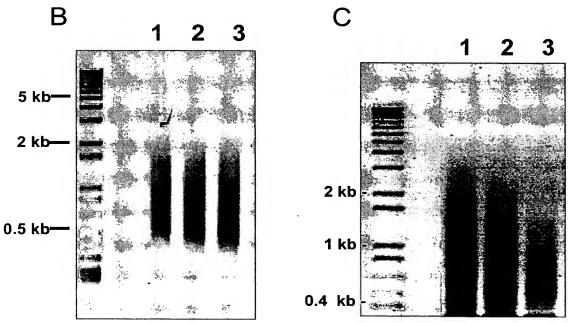


FIG. 15

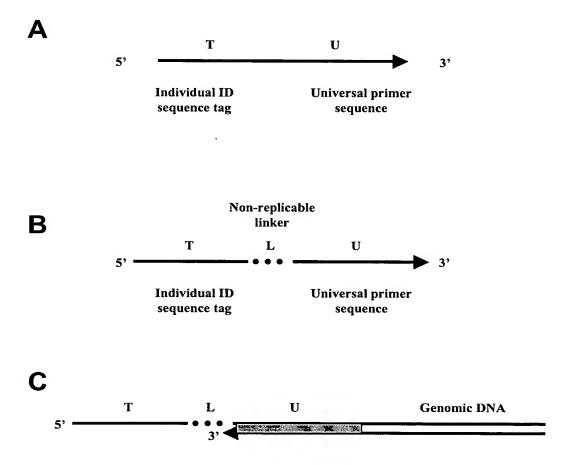


FIG. 16

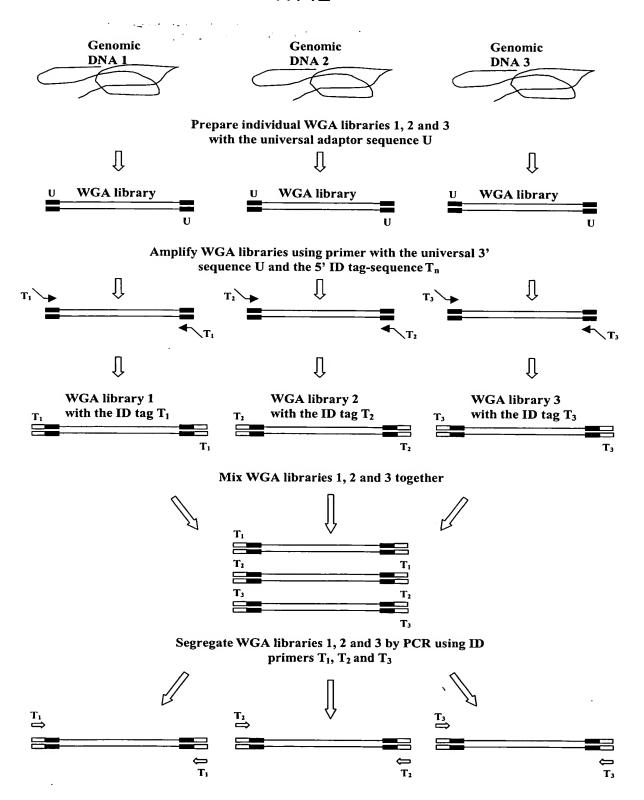
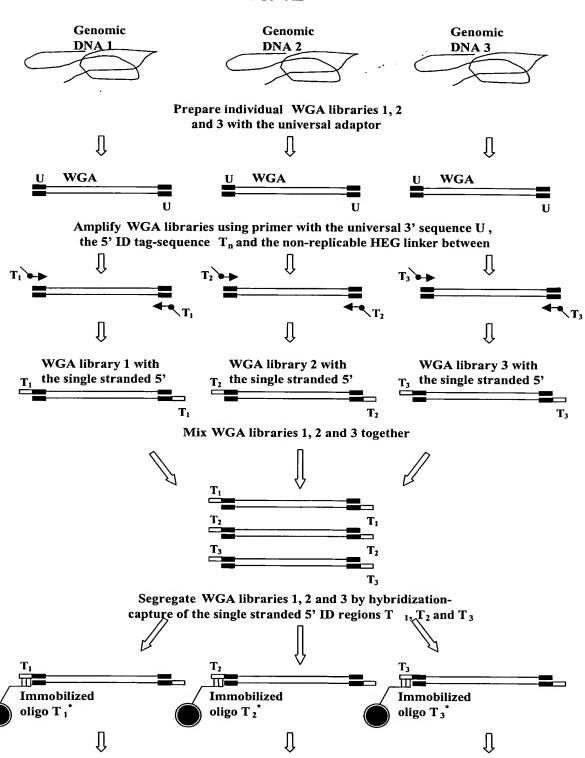
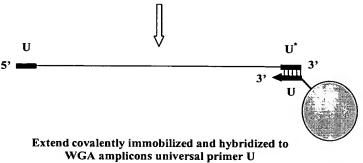


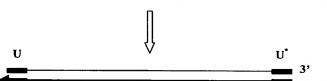
FIG. 17

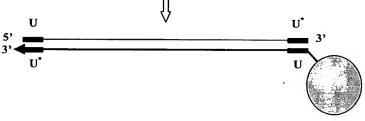


Amplify affinity-captured WGA libraries 1, 2 and 3 using universal primer U

FIG. 18







Wash with NaOH followed by TE

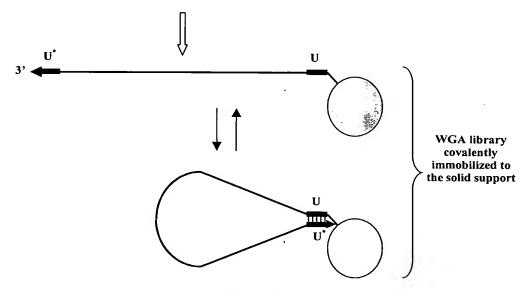
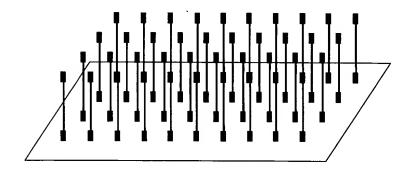
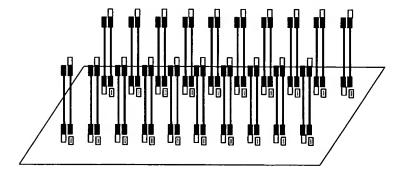


FIG. 19

A Covalent attachment of the individual WGA libraries to a WGA micro-array by replication

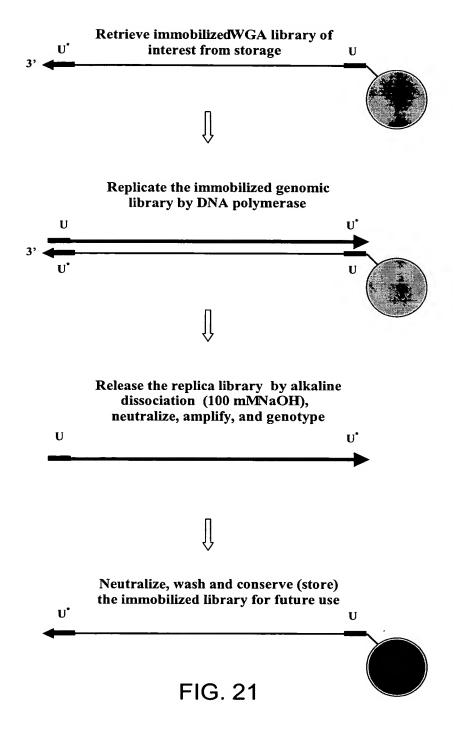


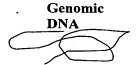
Non-covalent attachment of the individual WGA libraries to a WGA micro-array by hybridization



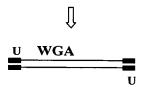
WGA libraries with the nonreplicable universal primer

FIG. 20

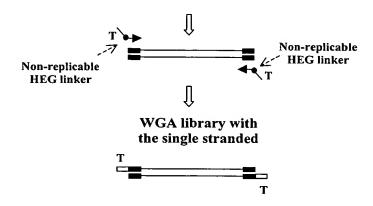




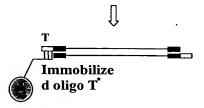
Prepare WGA library with the universal adaptor sequence U



Amplify WGA library using primer with the universal 3' sequence U, the 5' ID tag-sequence Tand the non-replicable HEG linker between them



Immobilize WGA library by hybridization-capture of the single stranded 5' IDregionT



Wash the immobilizee WGA library to remove contaminations

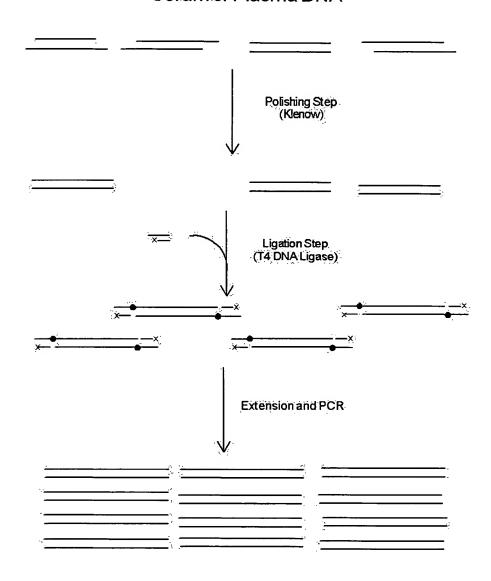


Release by heat

FIG. 22

A)

Serum or Plasma DNA

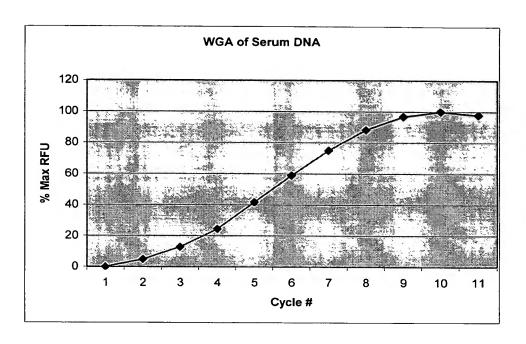


B)

Yb8 Forward: 5'-CGAGGCGGGTGGATCATGAGGT-3' (SEQ ID NO:48)

Yb8 Reverse: 5'-TCTGTCGCCCAGGCCGGACT-3' (SEQ ID NO:49)

A)



B)

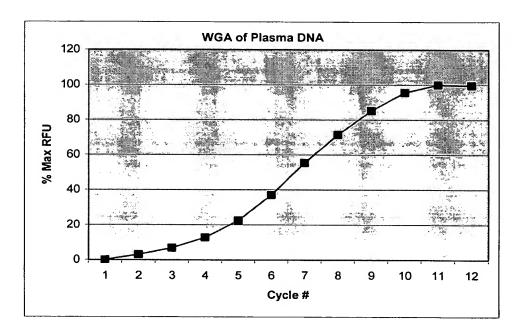
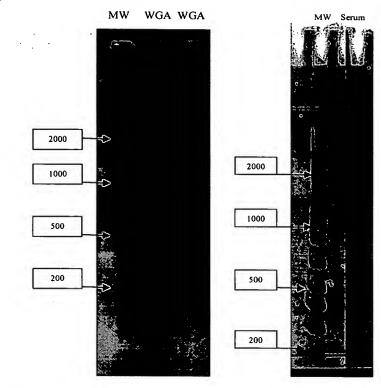


FIG. 24





B)

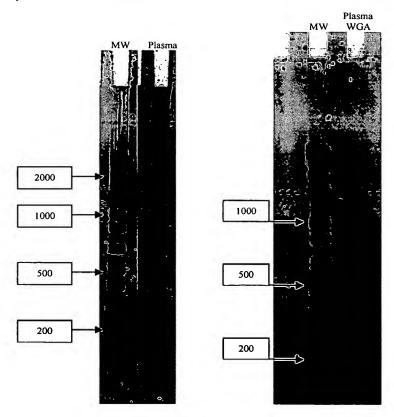


FIG. 25

STS Representation of Serum DNA and WGA Product

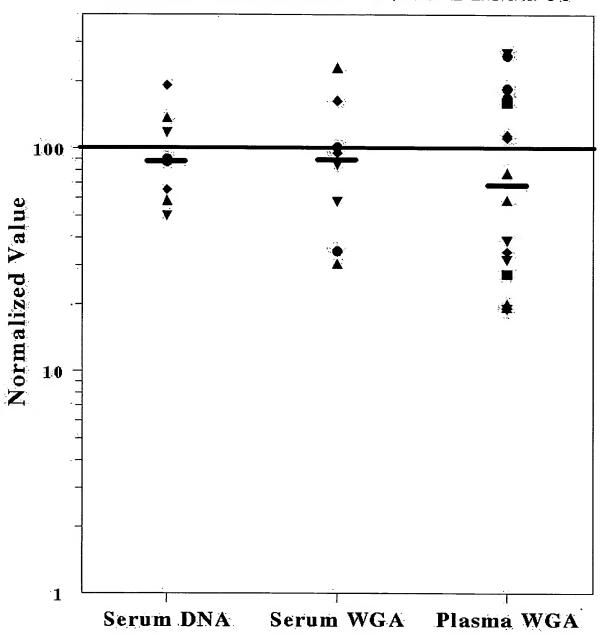


FIG. 26

Serum or Plasma DNA

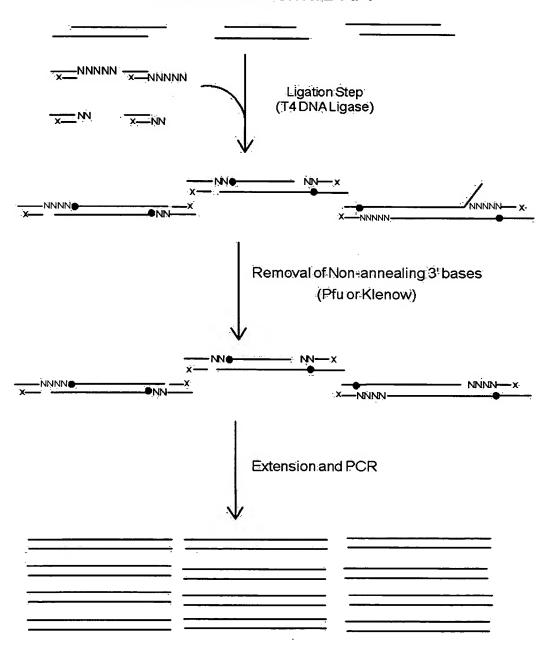


FIG. 27

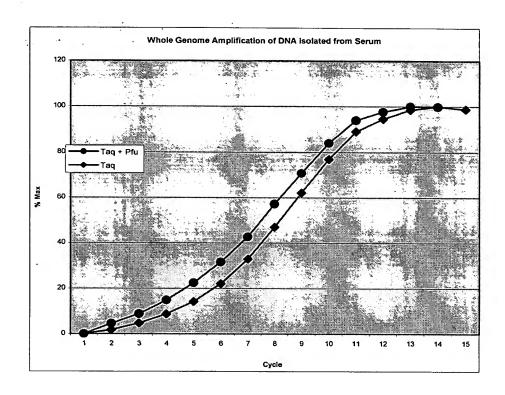
Figure Y2

5' T7N Overhang Adaptors

3' T7N Overhang Adaptors

T7N2 N2T7 (SEQ ID NO:44) 5'-GTAATACGACTCACTATAGGNN-3' 5-GTAATACGACTCACTATAGG-3' (SEQ ID NO:32) 3'-/3AmMC7/TGAGTGATATCC-5'
(SEQ ID NO:33) 3'-/3AmMC7/TGAGTGATATCCNN-5'(SEQ ID NO:12) **T7N3 N3T7** (SEQ ID NO:45) 5-GTAATACGACTCACTATAGG-3' (SEQ ID NO:32) 3'-/3AmMC7/TGAGTGATATCCNNN-5'(SEQ ID NO:13) 5'-GTAATACGACTCACTATAGGNNN-3' 3'-/3AmMC7/TGAGTGATATCC-5' (SEQ ID NO:33) **T7N4** N4T7 (SEQ ID NO:46) 5-GTAATACGACTCACTATAGG-3' (SEQ ID NO:32) 5'-GTAATACGACTCACTATAGGNNNN-3' 3'-/3AmMC7/TGAGTGATATCCNNNN-5' (SEQ ID NO:42) 3'-/3AmMC7/TGAGTGATATCC-5' (SEQ ID NO:33) **T7N5 N5T7** (SEQ ID NO:47) 5'-GTAATACGACTCACTATAGGNNNNN-3' 5-GTAATACGACTCACTATAGG-3' (SEQ ID NO:32) 3'-/3AmMC7/TGAGTGATATCC-5' 3'-/3AmMC7/TGAGTGATATCCNNNNN-5' (SEQ ID NO:43) (SEQ ID NO:33)

A)



B)

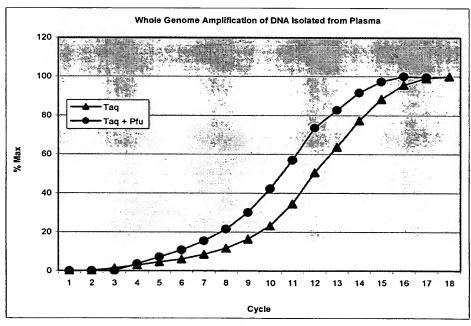
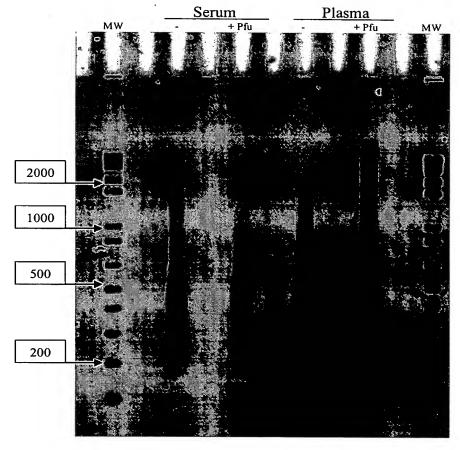


FIG. 29



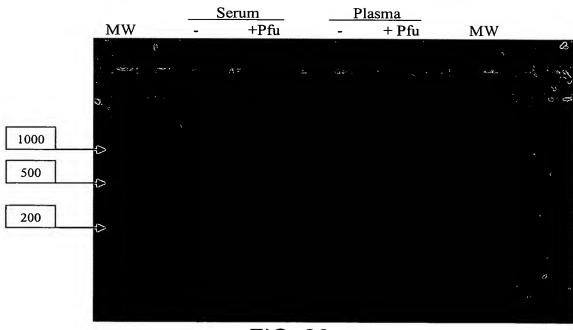


FIG. 30

STS Representation

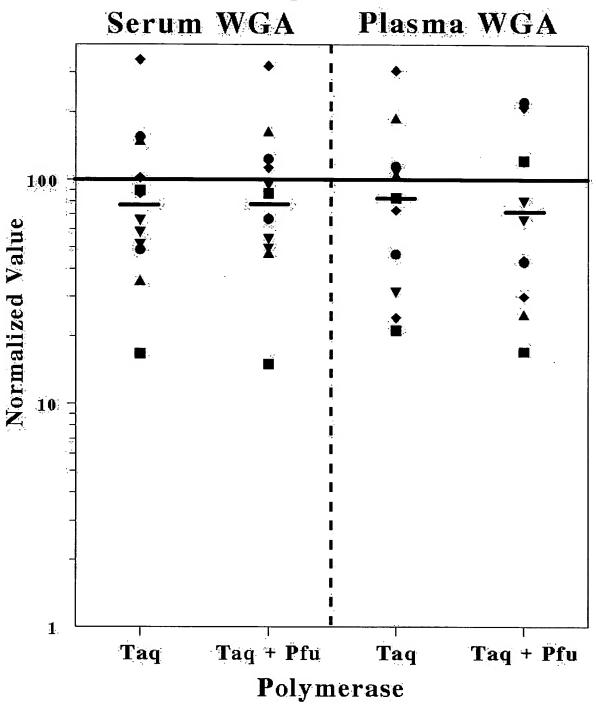


FIG. 31

Tissue

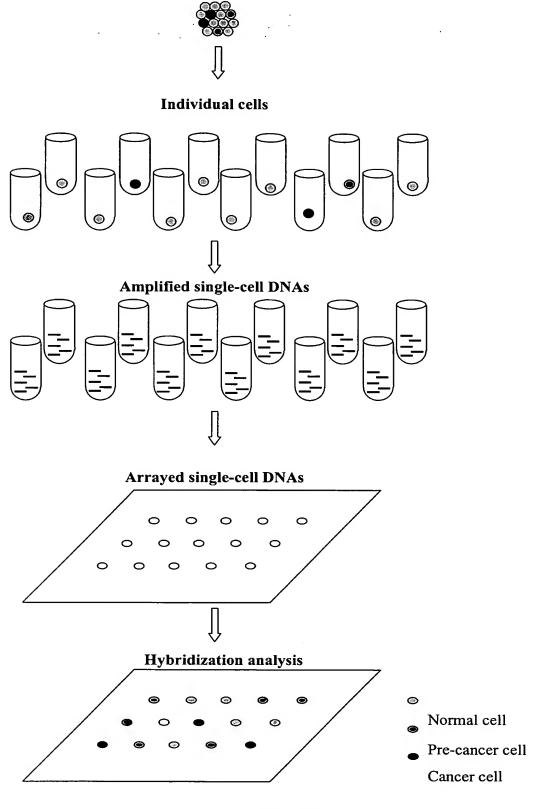


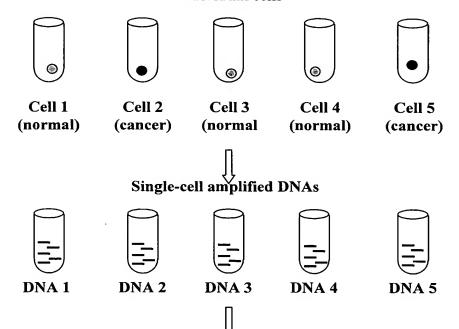
FIG. 32



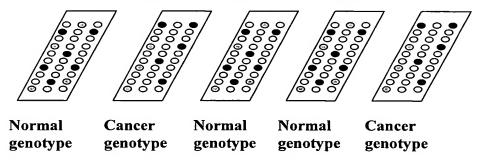




Individual cells



Tissue cell heterogeneity revealed by hybridization of the amplified single-cell DNAs to a micro-array



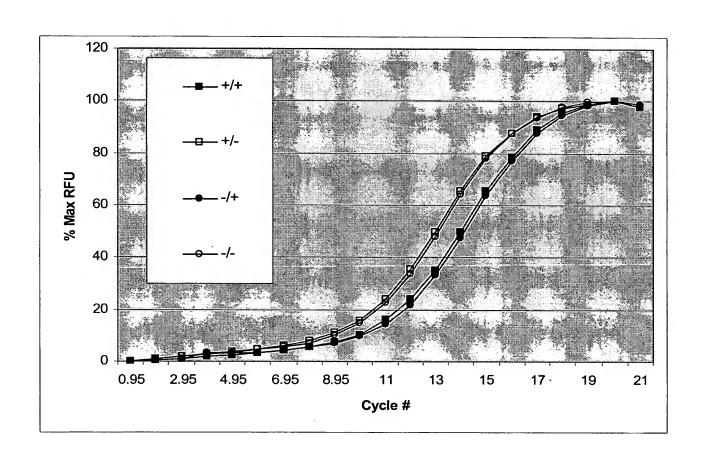
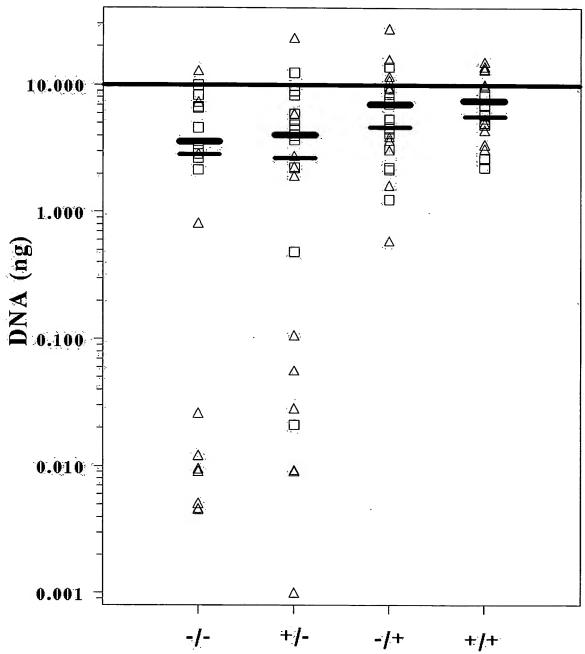


FIG. 34

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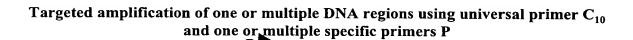
Representation of STS Sites



DMSO and N⁷-dGTP (Preparatin/Amplification)

FIG. 35

36/42 Prepare and amplify WGA or WTA library using universal primer sequence U U Add poly dG tail by Ligate adaptor Replicate amplified terminal containing C₁₀ WGA library using deoxynuclotidyl sequence at the 5' end modified primer U transferase of the long withC₁₀ sequence at oligonucleotide the 5' end G-tail C₁₀ G-tail



C₁₀

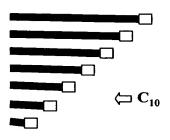
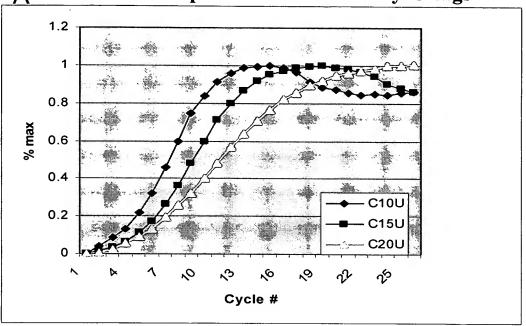
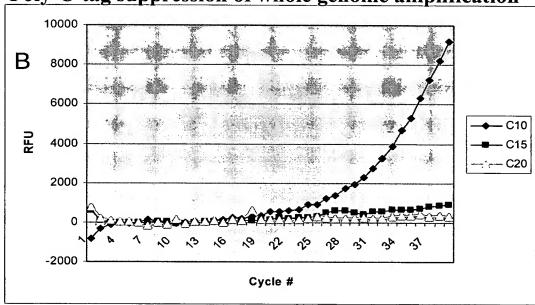


FIG.36

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WGA Incorporation terminal Poly-C-tags



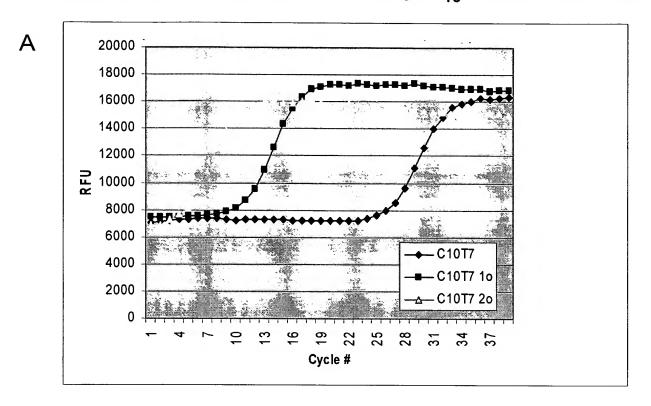
Poly-C-tag suppression of whole genome amplification



5'-CCCCCCCCC-3' (SEQ ID NO:53)
5'-CCCCCCCCCCCCCCC3' (SEQ ID NO:54)
5'-CCCCCCCCCCCCCCCCCC3' (SEQ ID NO:55)

FIG.37

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Targeted amplification from poly-C₁₀ WGA Libraries



TGA Specific Primer Titration

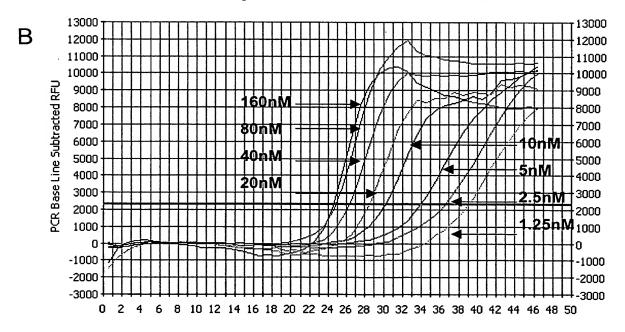
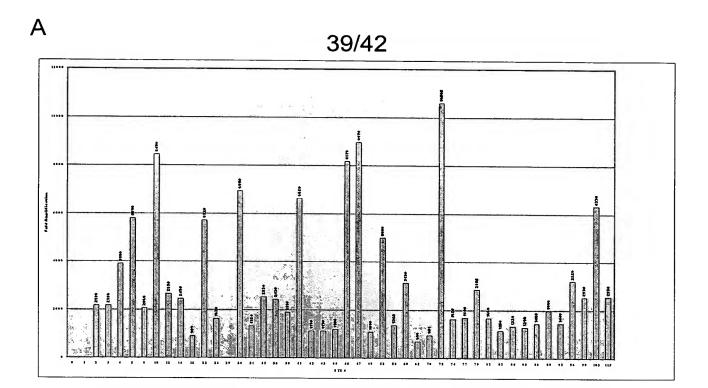


FIG. 38



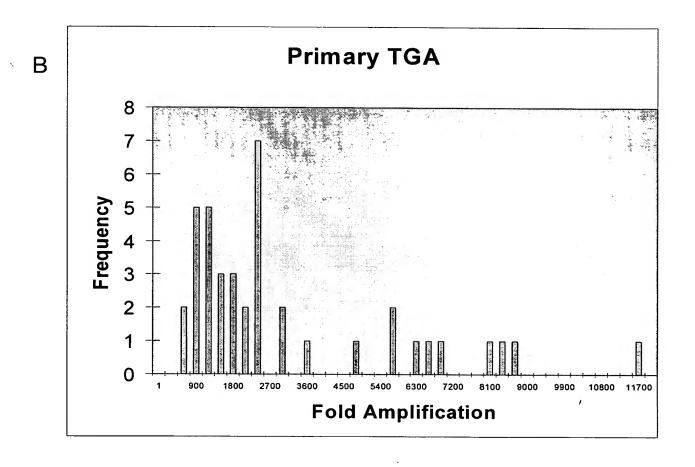
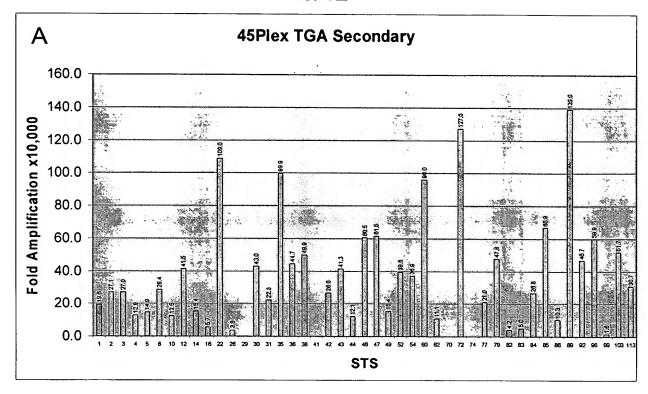


FIG. 39



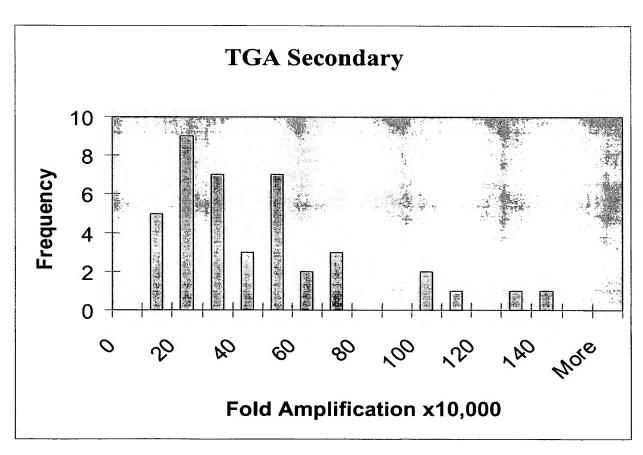


FIG. 40

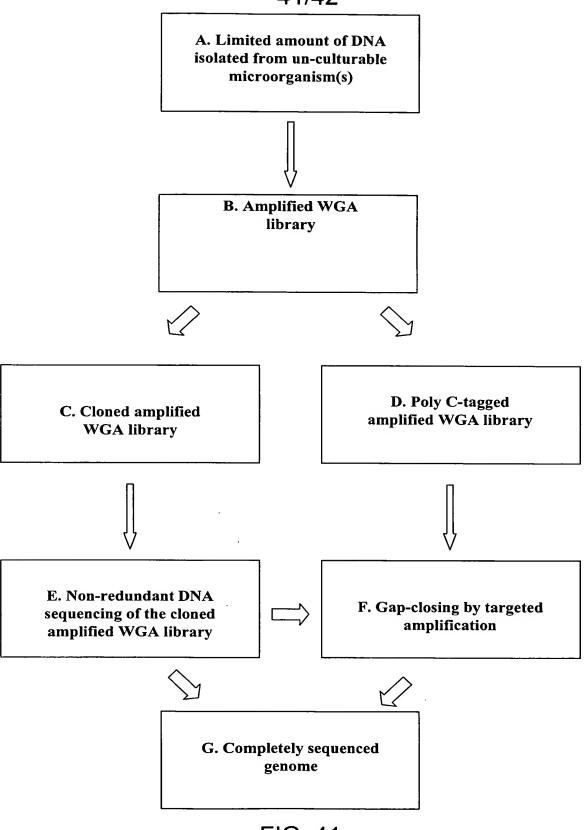


FIG. 41

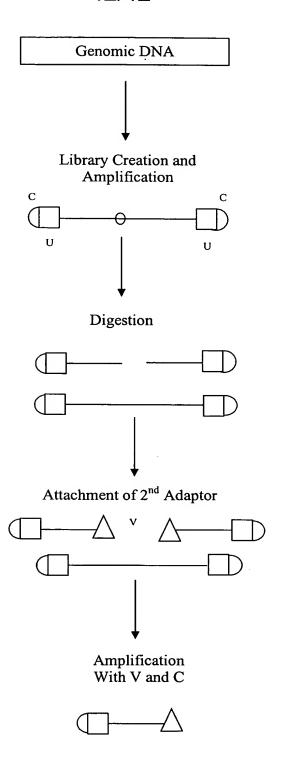


FIG. 42